



Substitute Form PTO-1449 (Modified)  <b>Information Disclosure Statement by Applicant</b> (Use several sheets if necessary)  (37 CFR §1.98(b))	U.S. Department of Commerce Patent and Trademark Office	Attorney's Docket No. <b>07064-010001</b>	Application No. <b>09/539,032</b>
	Applicant <b>Samir Kumar et al.</b>		
	Filing Date <b>March 30, 2000</b>	Group Art Unit <b>1431</b>	

**U.S. Patent Documents**

Examiner Initial	Desig. ID	Patent Number	Issue Date	Patentee	Class	Subclass	Filing Date If Appropriate
	AA						
	AB						
	AC						
	AD						
	AE						
	AF						
	AG						
	AH						
	AI						
	AJ						
	AK						

**Foreign Patent Documents or Published Foreign Patent Applications**

Examiner Initial	Desig. ID	Document Number	Publication Date	Country or Patent Office	Class	Subclass	Translation	
							Yes	No
	AL							
	AM							
	AN							
	AO							
	AP							

**Other Documents (include Author, Title, Date, and Place of Publication)**

Examiner Initial	Desig. ID	Document
<i>msc</i>	AQ	Altschul et al., "Basic Local Alignment Search Tool"; <u>J. Mol. Biol.</u> , Vol. 215, No. 3; pp 403-410 (1990)
<i>msc</i>	AR	Cutler et al., "TOR kinase homologs function in a signal transduction pathway that is conserved from yeast to mammals"; <u>Molecular and Cellular Endocrinology</u> , Vol. 155, Nos. 1-2; pp 135-142 (1999)
<i>msc</i>	AS	Ghannoum et al., "Antifungal Agents: Mode of Action, Mechanisms of Resistance, and Correlation of These Mechanisms with Bacterial Resistance"; <u>Clinical Microbiology Reviews</u> , Vol. 12, No. 4; pp 501-517 (1999)

Examiner Signature <i>My Moran</i>	Date Considered <i>8/24/02</i>
EXAMINER: Initials citation considered. Draw line through citation if not in conformance and not considered. Include copy of this form with next communication to applicant.	

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MS	AT	McCafferty et al., "Synergy and duality in peptide antibiotic mechanisms"; Chemical Biology, Vol. 3, No. 5; pp 672-680 (1999)
MS	AU	Porse et al., "Ribosomal Mechanics, Antibodies, and GTP Hydrolysis"; Cell, Vol. 97; pp 423-426 (1999)
MS	AV	Presnell et al., "A Segment-Based Approach to Protein Secondary Structure Prediction"; Biochemistry, Vol. 31, No. 4; pp 983-993 (1992)
MS	AW	Rooman et al., "Identification of predictive sequence motifs limited by protein structure data base size"; Nature, Vol. 335, No. 6185; pp 45-49 (1988)
MS	AX	Wilbur et al., "Rapid similarity searches of nucleic acid and protein data banks"; Proc. Natl. Acad. Sci USA, Vol. 80, No. 3; pp 726-730 (1983)
MS	AY	Wimberly et al., "A Detailed View of a Ribosomal Active Site: The Structure of the L11-RNA Complex"; Cell, Vol. 97, No. 4; pp 492-502 (1999)

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